\_\_\_\_\_\_

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Wed Jun 06 19:41:05 EDT 2007

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\*\*\*\*\*\*\*\*\*\*\*\*\*\*

Reviewer Comments:

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The above Xaa's are not explained; per 1.823 of the Sequence Rules, it is mandatory to explain all Xaa's (and n's, in a nucleotide sequence). Please indicate the Xaa locations on the <222> line, and indicate which amino acid residue(s) each Xaa represents on the <223> line. This error appears in subsequent sequences, too (e.g., 50-52, 54-55, etc.).

Also, the <213> mammalian response is invalid. Per 1.823 of the Sequence Rules, indicate Genus/species. This error appears throughout the sequence listing (e.g., Sequence 24, and many other sequences).

\*\*\*\*\*\*\*\*\*\*\*\*\*\*

## Validated By CRFValidator v 1.0.2

Application No: 10684346 Version No: 2.0

Input Set:

Output Set:

**Started:** 2007-06-05 17:52:42.790 **Finished:** 2007-06-05 17:52:44.912

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 122 ms

Total Warnings: 44

Total Errors: 0

No. of SeqIDs Defined: 167

Actual SeqID Count: 167

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Input Set:

Output Set:

**Started:** 2007-06-05 17:52:42.790 **Finished:** 2007-06-05 17:52:44.912

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 122 ms

Total Warnings: 44
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Actual SeqID Count: 167

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